

RAW SEQUENCE LISTING

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Application Serial Number: 10/516,813A
Source: IFWP
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PATENT APPLICATION: US/10/516,813A

DATE: 07/05/2006

TIME: 13:49:52

Input Set : A:\11752-007US1.txt

Output Set: N:\CRF4\07052006\J516813A.raw

5 <110> APPLICANT: FRASER, JOHN D.
 8 <120> TITLE OF INVENTION: IMMUNOMODULATORY CONSTRUCTS AND THEIR USES
 11 <130> FILE REFERENCE: 11752-007US1
 14 <140> CURRENT APPLICATION NUMBER: US 10/516,813A
 15 <141> CURRENT FILING DATE: 2004-12-03
 18 <150> PRIOR APPLICATION NUMBER: NZ 519371
 19 <151> PRIOR FILING DATE: 2002-06-04
 21 <150> PRIOR APPLICATION NUMBER: PCT/NZ03/00111
 22 <151> PRIOR FILING DATE: 2003-06-04
 25 <160> NUMBER OF SEQ ID NOS: 13
 28 <170> SOFTWARE: PatentIn version 3.1
 31 <210> SEQ ID NO: 1
 32 <211> LENGTH: 209
 33 <212> TYPE: PRT
 34 <213> ORGANISM: Streptococcus pyogenes
 37 <400> SEQUENCE: 1
 39 Leu Glu Val Asp Asn Asn Ser Leu Leu Arg Asn Ile Tyr Ser Thr Ile
 40 1 5 10 15
 43 Val Tyr Glu Tyr Ser Asp Ile Val Ile Asp Phe Lys Thr Ser His Asn
 44 20 25 30
 47 Leu Val Thr Lys Lys Leu Asp Val Arg Asp Ala Arg Asp Phe Phe Ile
 48 35 40 45
 51 Asn Ser Glu Met Asp Glu Tyr Ala Ala Asn Asp Phe Lys Thr Gly Asp
 52 50 55 60
 55 Lys Ile Ala Val Phe Ser Val Pro Phe Asp Trp Asn Tyr Leu Ser Lys
 56 65 70 75 80
 59 Gly Lys Val Thr Ala Tyr Thr Tyr Gly Gly Ile Thr Pro Tyr Gln Lys
 60 85 90 95
 63 Thr Ser Ile Pro Lys Asn Ile Pro Val Asn Leu Trp Ile Asn Gly Lys
 64 100 105 110
 67 Gln Ile Ser Val Pro Tyr Asn Glu Ile Ser Thr Asn Lys Thr Thr Val
 68 115 120 125
 71 Thr Ala Gln Glu Ile Asp Leu Lys Val Arg Lys Phe Leu Ile Ala Gln
 72 130 135 140
 75 His Gln Leu Tyr Ser Ser Gly Ser Ser Tyr Lys Ser Gly Arg Leu Val
 76 145 150 155 160
 79 Phe His Thr Asn Asp Asn Ser Asp Lys Tyr Ser Phe Asp Leu Phe Tyr
 80 165 170 175
 83 Val Gly Tyr Arg Asp Lys Glu Ser Ile Phe Lys Val Tyr Lys Asp Asn
 84 180 185 190
 87 Lys Ser Phe Asn Ile Asp Lys Ile Gly His Leu Asp Ile Glu Ile Asp
 88 195 200 205
 91 Ser

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95 <210> SEQ ID NO: 2
96 <211> LENGTH: 209
97 <212> TYPE: PRT
98 <213> ORGANISM: Streptococcus pyogenes
101 <400> SEQUENCE: 2
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104 1 5 10 15
107 Val Tyr Glu Tyr Ser Asp Ile Val Ile Asp Phe Lys Thr Ser His Cys
108 20 25 30
111 Leu Val Thr Lys Lys Leu Asp Val Arg Asp Ala Arg Asp Phe Phe Ile
112 35 40 45
115 Asn Ser Glu Met Asp Glu Tyr Ala Ala Asn Asp Phe Lys Thr Gly Asp
116 50 55 60
119 Lys Ile Ala Val Phe Ser Val Pro Phe Asp Trp Asn Tyr Leu Ser Lys
120 65 70 75 80
123 Gly Lys Val Thr Ala Tyr Thr Tyr Gly Gly Ile Thr Pro Tyr Gln Lys
124 85 90 95
127 Thr Ser Ile Pro Lys Asn Ile Pro Val Asn Leu Trp Ile Asn Gly Lys
128 100 105 110
131 Gln Ile Ser Val Pro Tyr Asn Glu Ile Ser Thr Asn Lys Thr Thr Val
132 115 120 125
135 Thr Ala Gln Glu Ile Asp Leu Lys Val Arg Lys Phe Leu Ile Ala Gln
136 130 135 140
139 His Gln Leu Tyr Ser Ser Gly Ser Ser Tyr Lys Ser Gly Arg Leu Val
140 145 150 155 160
143 Phe His Thr Asn Asp Asn Ser Asp Lys Tyr Ser Phe Asp Leu Leu Tyr
144 165 170 175
147 Val Gly Tyr Arg Asp Gln Glu Ser Ile Phe Lys Val Tyr Lys Asp Asn
148 180 185 190
151 Lys Ser Phe Asn Ile Asp Lys Ile Gly His Leu Asp Ile Glu Ile Asp
152 195 200 205
155 Ser
159 <210> SEQ ID NO: 3
160 <211> LENGTH: 209
161 <212> TYPE: PRT
162 <213> ORGANISM: Streptococcus pyogenes
165 <400> SEQUENCE: 3
167 Leu Glu Val Asp Asn Asn Ser Leu Leu Arg Asn Ile Tyr Ser Thr Ile
168 1 5 10 15
171 Val Ala Glu Tyr Ser Asp Ile Val Ile Asp Phe Lys Thr Ser His Cys
172 20 25 30
175 Leu Val Thr Lys Lys Leu Asp Val Arg Asp Ala Arg Asp Phe Phe Ile
176 35 40 45
179 Asn Ser Glu Met Asp Glu Tyr Ala Ala Asn Asp Phe Lys Thr Gly Asp
180 50 55 60
183 Lys Ile Ala Val Phe Ser Val Pro Phe Asp Trp Asn Tyr Leu Ser Lys
184 65 70 75 80
187 Gly Lys Val Thr Ala Tyr Thr Tyr Gly Gly Ile Thr Pro Tyr Gln Lys
188 85 90 95

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191 Thr Ser Ile Pro Lys Asn Ile Pro Val Asn Leu Trp Ile Asn Gly Lys
 192 100 105 110
 195 Gln Ile Ser Val Pro Tyr Asn Glu Ile Ser Thr Asn Lys Thr Thr Val
 196 115 120 125
 199 Thr Ala Gln Glu Ile Asp Leu Lys Val Arg Lys Phe Leu Ile Ala Gln
 200 130 135 140
 203 His Gln Leu Tyr Ser Ser Gly Ser Ser Tyr Lys Ser Gly Arg Leu Val
 204 145 150 155 160
 207 Phe His Thr Asn Asp Asn Ser Asp Lys Tyr Ser Phe Asp Leu Leu Tyr
 208 165 170 175
 211 Val Gly Tyr Arg Asp Gln Glu Ser Ile Phe Lys Val Tyr Lys Asp Asn
 212 180 185 190
 215 Lys Ser Phe Asn Ile Asp Lys Ile Gly His Leu Asp Ile Glu Ile Asp
 216 195 200 205
 219 Ser
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 224 <211> LENGTH: 27
 225 <212> TYPE: DNA
 226 <213> ORGANISM: Artificial sequence
 229 <220> FEATURE:
 230 <223> OTHER INFORMATION: Primer
 233 <400> SEQUENCE: 4
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 238 <211> LENGTH: 27
 239 <212> TYPE: DNA
 240 <213> ORGANISM: Artificial sequence
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 244 <223> OTHER INFORMATION: Primer
 246 <400> SEQUENCE: 5
 247 gaatattcag ctacaatcgt actata 27
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 251 <211> LENGTH: 24
 252 <212> TYPE: DNA
 253 <213> ORGANISM: Artificial Sequence
 256 <220> FEATURE:
 257 <223> OTHER INFORMATION: Primer
 259 <220> FEATURE:
 260 <223> OTHER INFORMATION: Primer
 262 <400> SEQUENCE: 6
 263 gatgttagat gtgctagaga ttcc 24
 266 <210> SEQ ID NO: 7
 267 <211> LENGTH: 26
 268 <212> TYPE: DNA
 269 <213> ORGANISM: Artificial sequence
 272 <220> FEATURE:
 273 <223> OTHER INFORMATION: Primer
 275 <400> SEQUENCE: 7
 276 ctctagcaca tctaacatca agttc 26

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279 <210> SEQ ID NO: 8
280 <211> LENGTH: 23
281 <212> TYPE: DNA
282 <213> ORGANISM: Artificial sequence
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289 ccatttgatt tgaactattt atc                                23
292 <210> SEQ ID NO: 9
293 <211> LENGTH: 23
294 <212> TYPE: DNA
295 <213> ORGANISM: Artificial sequence
298 <220> FEATURE:
299 <223> OTHER INFORMATION: Primer
301 <400> SEQUENCE: 9
302 gataaatagt tcaaataaaa tgg                                23
305 <210> SEQ ID NO: 10
306 <211> LENGTH: 23
307 <212> TYPE: DNA
308 <213> ORGANISM: Artificial sequence
311 <220> FEATURE:
312 <223> OTHER INFORMATION: Primer
314 <400> SEQUENCE: 10
315 gatatacgaga tcaagaaagt atc                                23
318 <210> SEQ ID NO: 11
319 <211> LENGTH: 23
320 <212> TYPE: DNA
321 <213> ORGANISM: Artificial sequence
324 <220> FEATURE:
325 <223> OTHER INFORMATION: Primer
327 <400> SEQUENCE: 11
328 gatactttct tgatctctat atc                                23
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332 <211> LENGTH: 18
333 <212> TYPE: DNA
334 <213> ORGANISM: Artificial sequence
337 <220> FEATURE:
338 <223> OTHER INFORMATION: Primer
340 <400> SEQUENCE: 12
341 accatcctcc aaaatcggt                                18
344 <210> SEQ ID NO: 13
345 <211> LENGTH: 18
346 <212> TYPE: DNA
347 <213> ORGANISM: Artificial sequence
350 <220> FEATURE:
351 <223> OTHER INFORMATION: Primer
353 <400> SEQUENCE: 13
354 tcagagggtt tcaccgtc                                18

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